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(MT)	

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MPsrch_pp protein - protein database search, using Smith-Waterman algorithm

Tabular output not generated. Run on: Sat May 13 08:52:23 2000; MasPar time 3.67 Seconds 212.807 Million cell updates/sec

Description:
Perfect Score:
Sequence: Title: >US-09-331-631-23 (1-33) from US09331631.pep 287 1 RSGRGECRRQCLRRHEGQPWETQECMRRCRRRG 33

Scoring table: PAM 150 Gap 11

188963 seqs, 23686106 residues

Post-processing: Minimum Match 0% Listing first 45 summaries

Searched:

Database: a-geneseq35 1:geneseqp

Statistics: Mean 23.079; Variance 98.498; scale 0.234

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result

esult No.	Score	Query Match	Length	DB	ID	Description	Pred. No.	
ш	287	100.0	33	-	W62836	Zea mays antimicrobial	3.17e-18	
2	287	00.	593	_	283	mays	.17e-1	
ω	269		35	-	R21079	0	.63e-1	
4	113	39.4	666	1	W62828	egrifoli	.95e-	
ر ت	109	•	625	ᆫ	W62830	integrifoli	٠	
σ	107		666	ᆫ	W62829		. 566	
7	88	30.7	590	_	W62832	Gossypium hirsutum ant		
∞	87	30.3	188	Ľ	W00726	መ	4.35e+00	
9	87	30.3	188	ш	W04829	coma vascular		
10	87	30.3	188	ш	W80493	Human vascular endothe	4.35e+00	
11	86	30.0		<u></u>	W62831	Theobroma cacao antimi		
12	86	30.0		μ,	R20181	Sequence encoded by 67		
13	84	29.3		۳	R28990		7.60e+00	
14	84	٠	215	μ	R28989	Encoded by GlcNAc rece	7.60e+00	
15	80	•	179	₽	W86221		1.58e+01	
16	80	٠	184	μ,	W86220	Human VEGF-3 truncated	1.58e+01	
17	80		189	_	W86219	Human VEGF-3 truncated	1.58e+01	
18	80	•	194	Ь	W86218	Human VEGF-3 truncated	1.58e+01	
19	80	•	201	_	W86236	Human VEGF-3 full leng	1:58e+01	
20	80	•	221	μ,	W07611	Human vascular endothe	1.58e+01	
21	80	•	221	щ	W86205	Human vascular endothe	1.58e+01	
22	80	27.9	388	Ц	Y04998	Mycobacterium species	1.58e+01	
23	76	26.5	110	-1	Y04944	Mycobacterium species	3.27e+01	

3.rag		and the second of the second o					Page
24	74	•	55	ᆸ	W80489	Murine vascular endoth	
25	74	25.8	55	Н	œ	Vascular endothelial g	4.68e+01
26	74	•	145	Н	W86213	Human VEGF-B truncated	
27	74	•	147	Н	W86212	VEGF-B	
28	74	•	150	Н	W86211	VEGF-B	
29	74	•	152	_	W86210	-B truncate	
30	74	•	155	ب	W86209	VEGF-B truncate	
31	74	•	160	ш	W86208	VEGF-B	
32	74	•	167	4سا	W86234	VEGF-B	
ω	74	•	188	 1	W00864	e VRF167.	
34	74	•	188	μ	W04826	Heart vascular endothe	
35	74	•	188	₽	W80490		
36	74	•	188	μ	W86201	ጠ	
37	74	•	195	Ļ	W04827	Heart vascular endothe	
38	74	•	195	μ	W80491	Murine vascular endoth	
39	72	•	125	Н	R28991	m.	
40	71	•	3025	Н	P93284	Sequence of clone HIV-	
<u>,,</u>	70	24.4	51	<u> </u>	W33694		
42	70	•	176	μ.	W33695		
43	70	•	387	<u> </u>	W18663	ented human N	
44	70	24.4	554	۰	W16873	1111111111111111111111	
45				۲	1140010	Bactitus churingiensis	9.51e+01

ALIGNMENTS

PT	PR RG	PN	X DE	D AC	RESULT	Qy	Дb	Qu Be Ma	SQ CC	888	PS.	ΡŢ	DR T	PA	מי א	PD	PN CS	X t		AC.	RESULT
77279/32. i-microbial protein from e.g. Macadamia in controlling microbial infestations of pl	02-JUL-1998. 22-DEC-1997; AU0874. 20-DEC-1996; AU-004275. (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY. ROWET NJ. GONTITER KC. GEOD II. Manners IM. Marches IP.	Zea mays. W09827805-A1:	anti bial	W62835; 27-OCT-1998 (first entry)	LT 2 W62835 standard; Protein; 593 AA.	1 RSGRGECHROCLERHEGOPWETOECMERCHERG 33	1 RSGRGECRRQCLERHEGQPWETQECMRRCRRG 33	Query Match 100.0%; Score 287; DB 1; Length 33; Best Local Similarity 100.0%; Pred. No. 3.17e-18; Matches 33; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	animais. Sequence 33 AA;	The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mammalian	tul for controlling microbial infestationsure; Page 60; 96pp; English.	al protein from e.g. Macadamia integr	Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP; wPI; 98-377279/32.	(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.	22-DEC-1997; AU0874. 20-DEC-1996: AU-004275	02-JUL-1998.	Zea mays. W09827805-A1.	antimicrobial protein; infestation; control.	1998 (first ent		W62836 standard. Drotein: 33 AA

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83228
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Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-JAN-1992.
05-JUN-1991; 305064.
05-JUN-1990; US-536127.
(PION-) PIONEER HI-BRED I
Duvick JP, Rood TA, Rao A
WPI; 92-010214/02.
                                                             22-DEC-1997; AU0874.
20-DEC-1996; AU-004275.
(RETR-) COOP RES CENT-TROPICAL PLANT PATHOLOGY.
Bower NI, Goulter KC, Green JL, Manners JM, Mar
WPI; 98-377279/32.
                                                                                                                                                                                                                                                                                                                                                     27-OCT-1998 (first entry)
Macadamia integrifolia antimicrobial
antimicrobial protein; infestation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Use of maize seed peptide CMIII and DNA encoding it - for killing or inhibiting plant pathogenic microorganisms.

Example 2; Page 5; 21pp; English.

Example 2; Page 5; 21pp; English.

The peptide (SEQ ID NO 1) was purified from public corn variety B73 and propriety corn variety MH18. It is basic and has has a total mol. wt. of 3900 daltons. The peptide sequence was used to design probes which were used to screen a maize genomic or cDNA.library. The isolated CMIII gene can be used to prepare an expression vector for prodn. of recombinant CMIII for use in controlling plant patho-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             be used animals.
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The se
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         useful
                           Novel anti-microbial protein
                                               N-PSDB; V42310
                                                                                                                                                                                                                                                                                            Peptide
                                                                                                                                                                                                                                                                                                                                       Macadamia
                                                                                                                                                                                                                                                                                                                                                                                                                           W62828 standard;
W62828;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zea mays.
EP-465009-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ic organisms.
also Q20272 and 3.
uence 35 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      im 1; Page 58-60; 96pp;
sequence is that of an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RSGRGECRRQCLRRHEGQPWETQECMRRCRRRG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
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    for controlling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
Similarity 100.0%;
33; Conservati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        h 93.7%;
Similarity 97.0%;
32; Conservative
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ce is that of an antimicrobial protein which can
control microbial infestations in plants and mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        593 AA;
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corn; pathogen
                                                                                                                                                                                                                                          /note- "signal peptide" 29. .666
                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide;
  rotein from e.g. Macadamia integrifolia microbial infestations of plants or man
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Pred. No. 1.63e-16;
0; Mismatches 1;
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Pred.
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No. 3.17e-18;
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                                                                                          JM, Marcus
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    or mammals
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85008
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Best Local S
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                      WO982780.5-A1.
02-UTL-1998;
22-DEC-1997; AU0874.
20-DEC-1996; AU-004275.
(RETR-) COOP RES CENT TR
BOWER NI, Goulter KC, Gr
WPI; 98-377279/32.
N-PSDB; V42311.
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20-DEC-1996; AU-004275.
(RETR-) COOP RES CENT TR
BOWER NI, GOULLER KC, Gr
WPI; 98-377279/32.
N-PSDB; V42316.
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animals.
Sequence
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Macadamia integrifolia antimicrobial
antimicrobial protein; infestation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 43-45; 96pp; English.
The sequence is that of an antimicrobial be used to control microbial infestations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel anti-microbial protein useful for controlling microbial for controlling microbial in page 43-45; 96pp; En The sequence is that of an an
                                                                                                                                                                                                                                                                                          Key
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    Novel anti-microbial
                                                                                                                                                                                                                        Protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6 ECRRQCLRRHEGQPWETQECMRRCR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QCQRRC-RQQESDPRQQQYCQRRCK
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                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               h 38.0%;
Similarity 44.0%;
11; Conservative
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protein; infestation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          625 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        666 AA;
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is that of an
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                                                                                                                                                                                                 29. .60
/note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "signal peptide" 29. .666
                                                                                                                                                                                                                                          note= "signal"
                                                                                                                                                                                                                                                                                   ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                       Protein;
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                                                                 Green JL,
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antimicrobial protein which can
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Pred. No. 2.9
7; Mismatch
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Pred. No. 6.47e-02;
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RESULT RESULT AND ACCOUNT OF A COUNTY AND ACCOUNTY AND AC
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                                                                                                                      Weber
WPI; 9
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22-DEC-1996; AU-004275.
20-DEC-1996; AU-004275.
(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
BOWER NI, Goulter KC, Green JL, Manners JM, Mar
WPI; 98-377279/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  useful for controlling microbial infestations of plants or mammals Claim 1; Page 39-41; 96pp; English.
The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mammalian animals.
                                                                                                                                                                                    22-FEB-1996; AU0094.
02-MAR-1995; AU-001457.
20-NOV-1995; AU-006647.
22-DEC-1995; AU-007274.
(AMRA-) AMRAD OPERATIONS PTY LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel anti-microbial protein from e.g. Macadamia integrifolia - useful for controlling microbial infestations of plants or mammals Claim 1; Page 49-51; 96pp; English.

The sequence is that of an antimicrobial protein which can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        W62832 standard; Protein; 590
W62832;
27-OCT-1998 (first entry)
                                                                     N-PSDB; T33611.
New growth factor related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vascular endothelial growth Vascular endothelial growth
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W00726 standard;
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                                            useful for inducing astroglial
                                                                                                                                                                Grimmond S,
                                                                                                                                                                                                                                                                                                           06-SEP-1996.
                                                                                                                                                                                                                                                                                                                                    WO9627007-A1.
                                                                                                                                                                                                                                                                                                                                                                                peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                astroglial proliferation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      W00726;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     128 QCQERC-QRHETEPRHMQTCQQRCERR 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
les 12; Conser
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    12;
Page 42-43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 590 AA;
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                                                                                                                                                                   Hayward NK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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113pp;
                                                                                                                                                                                                                                                                                                                                                       Sig_peptide
                                                                       to vascular endothelial growth factor
                                                                                                                                                                   Larsson C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antimicrobial protein which
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Pred. No. 9.
7; Mismatc
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factor; VEGF; SOM175-e6; neuron;
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English
                                            proliferation
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9.56e-02;
7;
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                                              and promoting neuronal
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TRESULT PROPERTY OF THE PROPER
                                                                                                                         CC W04824-W04831 represent the vascular endothelial growth factor (VEGF) CC proteins of the invention, which promote endothelial or mesodermal cell CC proliferation. VEGF is also a glycosylated cationic dimer, and is CC sometimes referred to as vascular permeability factor (VPF). VEGF has CC diverse effects, depending on the specific biological context in which it is found. VEGF is a potent endothelial cell mitogen, and directly CC contributes to induction of angiogenesis in vivo by promoting endothelial cell growth during normal embryonic development, wound healing, and CC tissue regeneration/reorganisation. The VEGF proteins of the invention CC share the angiogenic and other properties of VEGF, but are distributed and expressed in tissues differently to VEGF. The proteins can therefore be used to accelerate angiogenesis in wound healing. Antibodies against the proteins can be used for inhibiting angiogenesis. The antibodies can complementary to the coding sequences for the proteins of the invention CC analso be used to detect VEGF-B coding sequences. Quantification of VEGF-B in cancer biopsy specimens may be useful as an indicator of metastatic risk. VEGF-B expression in a cell can be retarded using sequence 1988 AA; in retarding VEGF expression in tumour cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  888888888888
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Best Local Similarity
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06-JUN-1995; US-469427.
06-DEC-1995; US-469427.
06-DEC-1995; US-569063.
(LUDW-) LUDWIG INST CANCER RES.
(LUDW-) LUDWIG INST CANCER RES.
(UYHE-) UNIV HELSINKI LICENSING LTD OY
Alitalo K, Eriksson U, Olofsson B, I
WPI; 96-412582/41.
N-PSDB; T37913.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 18; Page 59; 107pp; English. W04824-W04831 represent the protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fibrosarcoma vascular endothelial growth factor-B167. Endothelial cell; proliferation; vascular endothelial growth factor; VPF Endothelium; mesodermal cell; cationic dimer; tissue regeneration; vascular permeability factor; cell mitogen; angiogenesis; cell growth; embryonic development; wound healing; tissue reorganisation; antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Splice variants (W00726-28) of the human vascular endothelial growth factor-like polypeptide SOM175 (see also W00725) are products of cDNA clones (see also T33611-13) respectively lacking exon 6, exons 6+7, and exon 4 of the SOM175 gene (see also T33610). They show at least 1 of the properties of SOM175 including the ability to induce proliferation of vascular endothelial cells, to interact with flt-J/flk-1 receptors, and to induce cell migration, cell survival and/or an increase in intracellular levels of alkaline phosphatase. Recombinant SOM175 proteins can be used to induce astrogical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9626736-A1.
06-SEP-1996.
01-MAR-1996; U02957
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      30.3%;
larity 40.6%;
Conservative
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Pred.
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Pred. No. 4.35e+00;
7; Mismatches 9
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e 87; DB 1; Leny
.. No. 4.35e+00;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      or VEGF-B proteins - useful to healing, also related nucleic
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B, Pajusola F
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                                                                   Length 188;
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01-MAR-1996; 05-609443.

01-MAR-1996; US-609443.

01-MAR-1995; US-397651.

06-JUN-1995; US-469427.

06-DEC-1995; US-569063.

(LUDW-) LUDWIG INST CANCER RES.

(UYHE-) UNIV HELSINKI LICENSING LTD OY.

Alitalo K, Eriksson U, Olofsson B, Pajusc

WPI; 99-034079/03.
                                                                                                                                                                             02-JUL-1998.
22-DEC-1997; AU0874.
20-DEC-1996; AU-004275.
(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
Bower NI, Goulter KC, Green JL, Manners JM, Marv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents human vascular endothelial growth factor (VEGF)-B167. VEGF proteins are used for promoting proliferation of endothelial cells and for stimulating angiogenesis (the proliferation of new capillaries form pre-existing blood vessels). These activities are useful for treating tissue growth and repair, including organ repair. This is also useful in pregnancy, in follicle development, as these processes must occur in development of the placenta. The proteins can also be used to raise antibodies, either for use in detection of the proteins or as inhibitors of their action. This is especially useful as angiogenesis is required by tumours as they need new blood supplies to grow and proliferate.
animals.
Sequence
                                                                         Novel anti-microbial protein from e.g. Macadamia useful for controlling microbial infestations of Claim 1; Page 47-49; 96pp; English.
The sequence is that of an antimicrobial protein
                                                      be used to control microbial infestations
                                                                                                                                                                                                                                                                                                                                                          Theobroma cacao. W09827805-A1.
                                                                                                                                                                                                                                                                                                                                                                                                        Theobroma cacao antimicrobial protein. antimicrobial protein; infestation; control.
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27-OCT-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Vascular endothelial growth factor-B iso:forms, and DNA encoding them useful for inducing anglogenesis and cellular proliferati and raising antibodies to inhibit activities in e.g. tumours Claim 1; Fig 11; 52pp; English.
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Homo sapiens.
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W80493;
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Similarity 40.6%;
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growth factor; VEGF; proliferation;
                                                antimicrobial protein which can
al infestations in plants and mammalian
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7; M
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      87; DB 1; L
No. 4.35e+00;
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Spencer ME, Hodge R, Deakin EA, Ashton WPI; 92-024418/03.
N-SDB; Q20377.
or diagnostic use Disclosure; Fig 5a: 75pp; French.
Clone'TGR-CLIbis (see Q31005) was isolated by immunoscreening Western blot derived from a cDNA bank of normal human thyroid in lambda gtl. Denatured GlCNAc receptor was purified from pothyroids and used to raise polyclonal antibodies in rabbits for immunoscreening. Sequence TGR-CLI is complementary to
                                                                                                                                    30-APR-1992; F00396.

03-MAY-1991; FR-005478.

(CNRS) CENT NAT RECH SCI.

Blanck O, Courageot J, Miquelis

WPI: 92-39886/48.
                                                                                                                                                                                                                                            Encoded by clone TGR-CL1 complementary to TGR-CL1bis. Thyroid N-acetyl-glucosamine receptor; carbohydrate recoglcnAc receptor; CRD; thyroid adenocarcinoma; lysosome; Hashimoto's disease; Basedow's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The inventors claim a 67 kD and 31 kD T. cacao protein, and fragments, and encoding DNAs. The 47 kD and 31 kD proteins are derived from the 67 kD precursor. T. cacao protein cDNA was detected in a cDNA library prepared from immature cocca beans RNA using a probe based on the AA sequence of a CNBr peptide common to the 47 kD and 31 kD polypeptides. Homology searches revealed close homologies between the 67 kD polypeptide and the vicilins, which are
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                                                                                            New thyroid N-acetyl-glucosamine receptor proteins fragments, antibodies and encoding nucleic acids, i
                                                                                                                          N-PSDB; Q31014
                                                                                                                                                                                                                        W09219733-A.
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06-APR-1993 (first entry)
                                                                                                                                                                                                                                                                                                                                   R28990 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     beans and produced expression vectors
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07-JUN-1991; G00914
11-JUN-1990; GB-013
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                                                                                                                                                                                                                                      Homo sapiens:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 4; Fig 2; 59pp; The inventors claim a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Recombinant cocoa proteins -
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ence 566 AA;
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larity 36.0%;
Conservative
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oded by 67 kD T.
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-glucosamine receptor; carbohydrate recognition domain;
                                                                                                                                                                                                                                                                                                                                    Protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                proteins - are responsible for flavour in cocoa in large quantities using yeast and bacterial
                                                                                                                                                                                                                                                                                                                                                                                                                                                          30.0%;
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67 kD and
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Pred. No. 5
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RESULT ID WE AC DT 11 DE HU DE HU DE KWW C C C KWW C C KWW C C KWW C KWW C KWW C C
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                          16-FEB-1999 (first entry)
16-FEB-1999 (first entry)
Human VEGF-3 truncated fragment 4.
VEGF: VRP; vascular endothelial growth factor; VEGF-related protein; recombinant; truncated; gene therapy; angiogenesis; cardiac ischaemia; coronary; collateral vessel development; cell growth; migration; heart; lower limb ischaemia; stroke; peripheral vascular disease; intestine; wound healing; skin; vascular permeability.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                respectively) were isolated by immunoscreening a Western blot derived from a cDNA bank of normal human thyroid cDNA in lambda gtll. Denatured GlcNAc receptor was purified from porcine thyroids and used to raise polyclonal antibodies in rabbits for use in immunoscreening. Sequences TGR-CL1 and TGR-CL5 are complementary to the corresponding "bis" sequences and partially overlap one another. The amino acid sequence deduced from the combined overlapping sequences is not a GlcNAc receptor. See Q31005-Q31018.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-APR-1992; F00396.
03-MAX'1991; FR-005478.
(CNRS ) CENT NAT RECH SCI.
Blanck O, Courageot J, Miquelis R, Thibault V;
WPI; 92-398860/48.
N-PSDB; R28989.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Encoded by GlcNAc receptor complementary sequence. Thyroid N-acetyl-glucosamine receptor; carbohydrate recognition domain; Thyroid nearcarcinoma; lysosome; endosome; TGR-CL1; TGR-CL5; Hashimoto's disease; Basedow's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the corresponding "bis" sequence and includes an sequence deduced from the complementary sequence receptor. See Q31005-Q31018.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or diagnostic use
Disclosure; Fig 3; 75pp; French.
Clones TGR-CLIbis and TGR-CL5bis (see Q31005 and Q31007,
     Homo sapiens
WO9849300-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New thyroid N-acetyl-glucosamine receptor proteins - and fragments, antibodies and encoding nucleic acids, for therapeutic
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                                                                                                                                                                                                                                                                                     W86221 standard; protein; 179
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                                                                                                                                                                                                                                                                                                                                                                                                                                             108 GRSPCRHPCAPRQSQCAPWSGP-CAQ-CWR 135
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nes 11; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27 GRSPCRHPCAPRQSQCAPWSGP-CAQ-CWR 54
                                                                                                                                                                                                                                                                                                                                                                                                         3 GRGECRRQCLRRH-EGQPWETQECMRRCRR 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ω
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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/note= "overlap"
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36.7%;
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36.7%;
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Pred. No. 7.60e+00;
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Pred. No. 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6
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7.60e+00;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 215;
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Best Local Similarity
Matches 13; Conser
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20-APR-1998; U7801.
25-APR-1997; US-842984.
(COLL-) COLLATERAL THERA
Bohlen P;
WPI; 99-009426/01.
                                                                                                                                                                                                                                                                       The invention relates to truncated VRP (vascular endothelial growth factor (VEGF)-related protein) subunits that have at least one amino acid N-terminal to the first Cys of the core sequence deleted. Host cells transformed or transfected with expression vectors containing nucleic acids encoding the truncated VRP subunits are used to produce the truncated proteins recombinantly. The truncated VRP subunits, optionally expressed from gene therapy vectors, have in vivo and in vitro angiogenic activity and are used to stimulate angiogenesis, particularly coronary collateral vessel development in cases of cardiac ischaemia; to stimulate endothelial cell growth and migration in vitro; to treat heart disease; to treat ischaemia (e.g.cardiac, chronic coronary or chronic lower limb ischaemia; stroke and peripheral vascular disease); to promote healing of wounds (of skin or intestines), and to increase vascular permeability.

Sequences W86218 to W86221 represent truncated fragments of VEGF-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New truncated vascular endothelial growth factor-related protein subunits - lack part of the N-terminal sequence, used to stimulate angiogenesis, e.g. for treating heart disease and ischaemia claim 5; Fig 2C; 113pp; English
                                                                                                                                                                                                                                                            Sequence
                                                                                                                              128
                                                                                     2
                                                                                                              SPRPLCPR-CTQHHQC-P-DPRTCRCRCRRRS 156
                                                                                   SGRGECRRQCLRRHEGQPWETQECMRRCRRRG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COLLATERAL THERAPEUTICS
                                                                                                                                                                                                                                                            179 AA;
                                                                                                                                                                      Conservative
                                                                                                                                                                                             27.98;
                                                                                                                                                                      Pred. No. 1.58e+6; Mismatches
                                                                                                                                                                                             Score 80; DB 1;
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